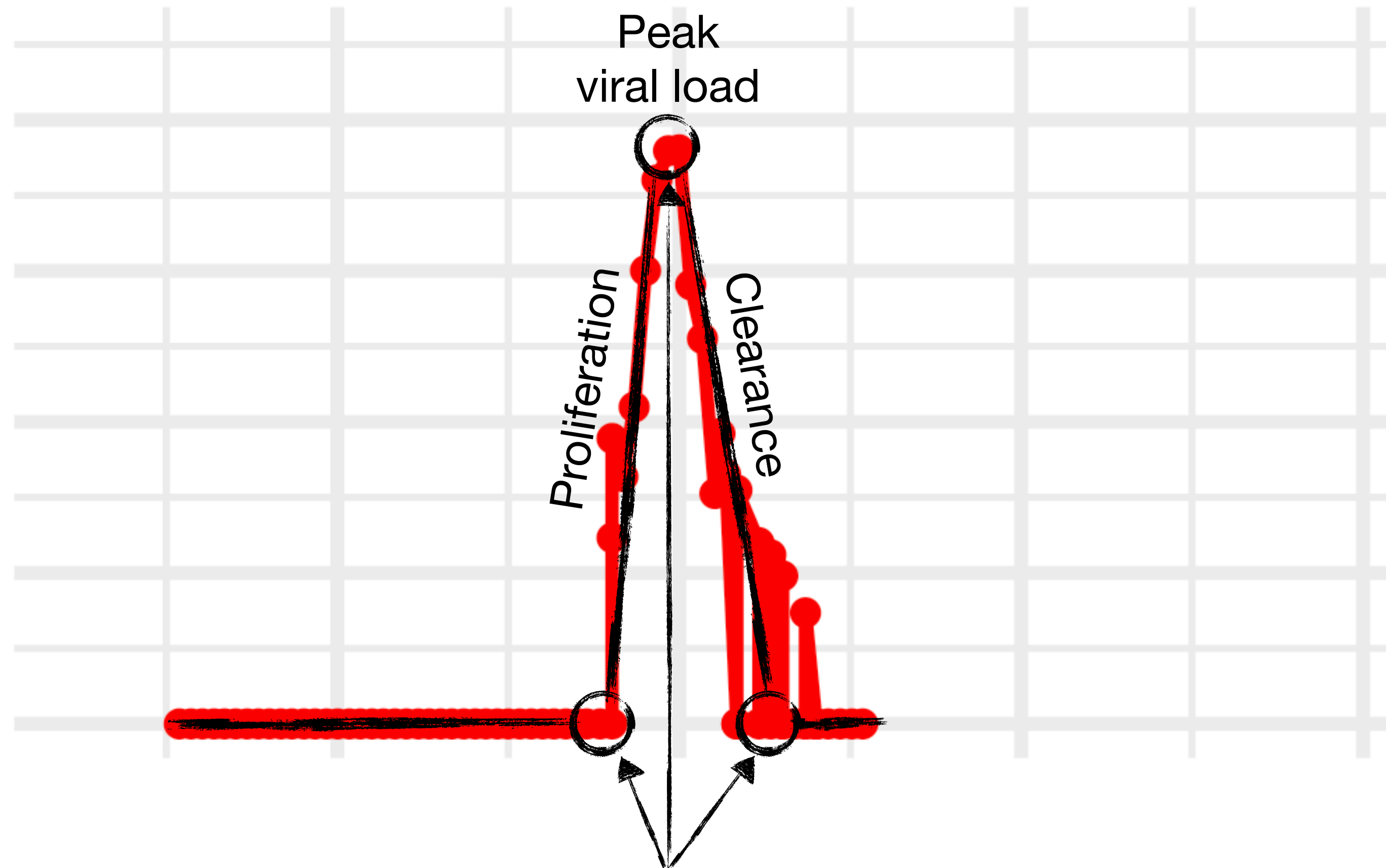


# A model for SARS-CoV-2 viral kinetics

*Ça monte, ça descend:*



**Goal:** estimate where these “hinge points” should lie to optimize the fit to the data

# A model for SARS-CoV-2 viral kinetics

*Ça monte, ça descend:*

